

MSA length = 2297

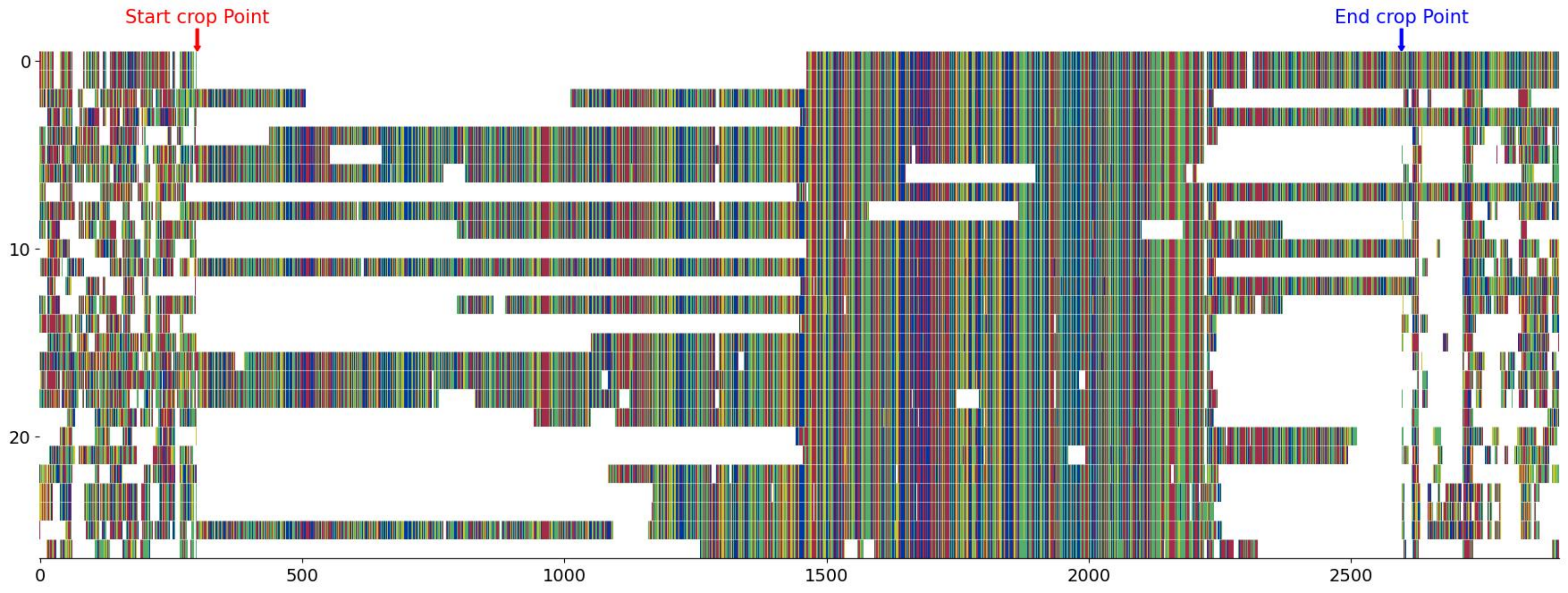
Start crop Point

End crop Point



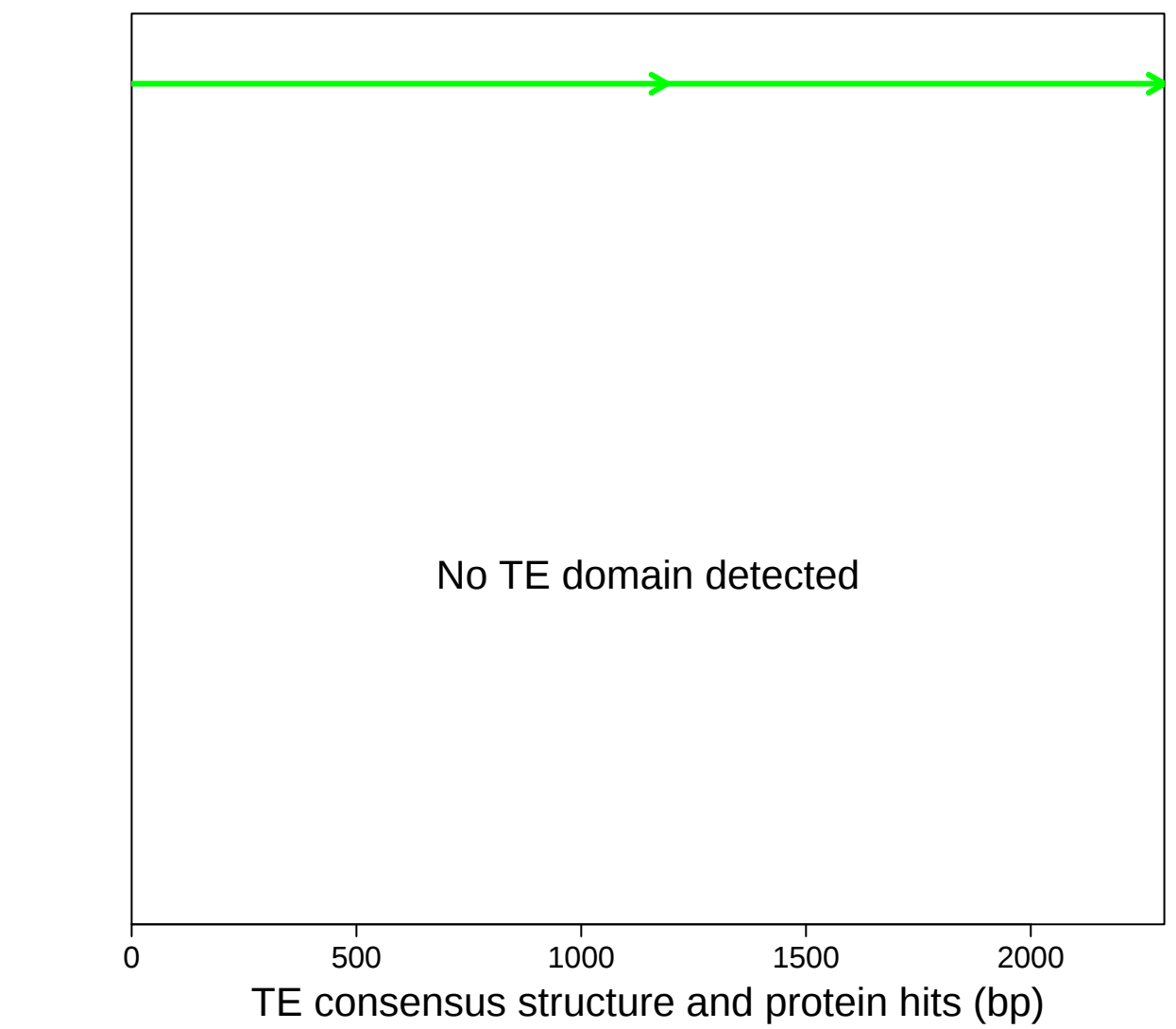
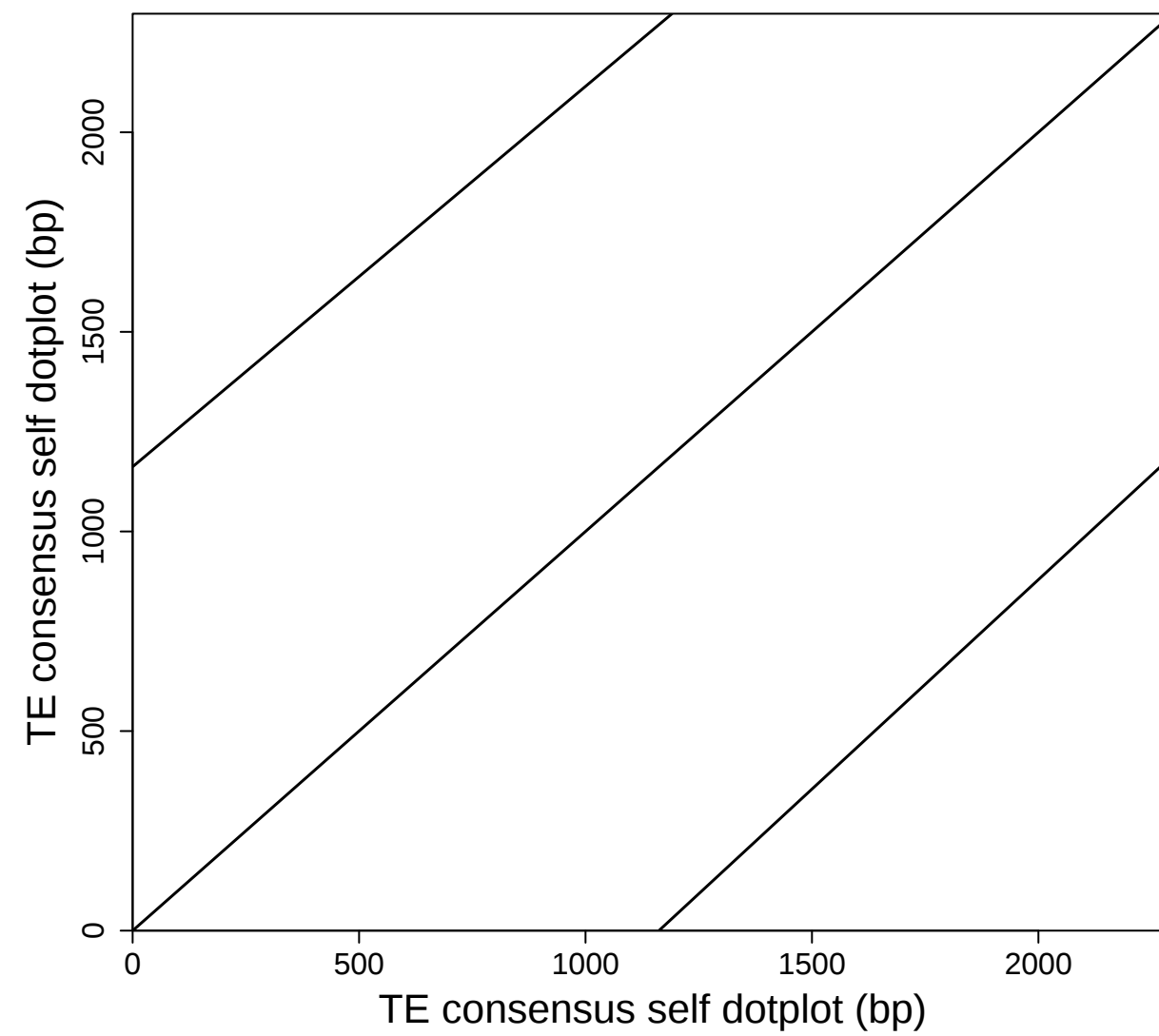
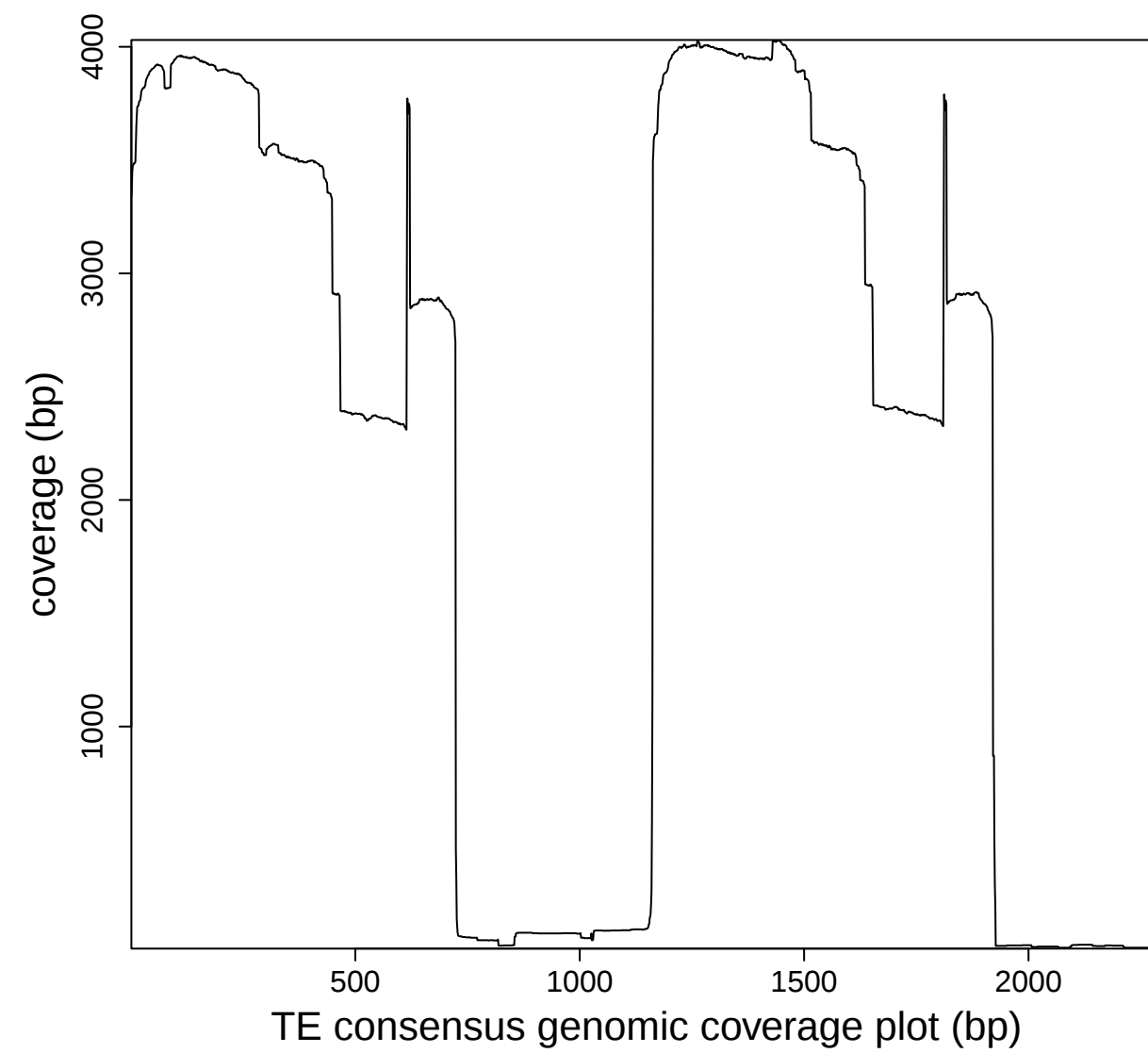
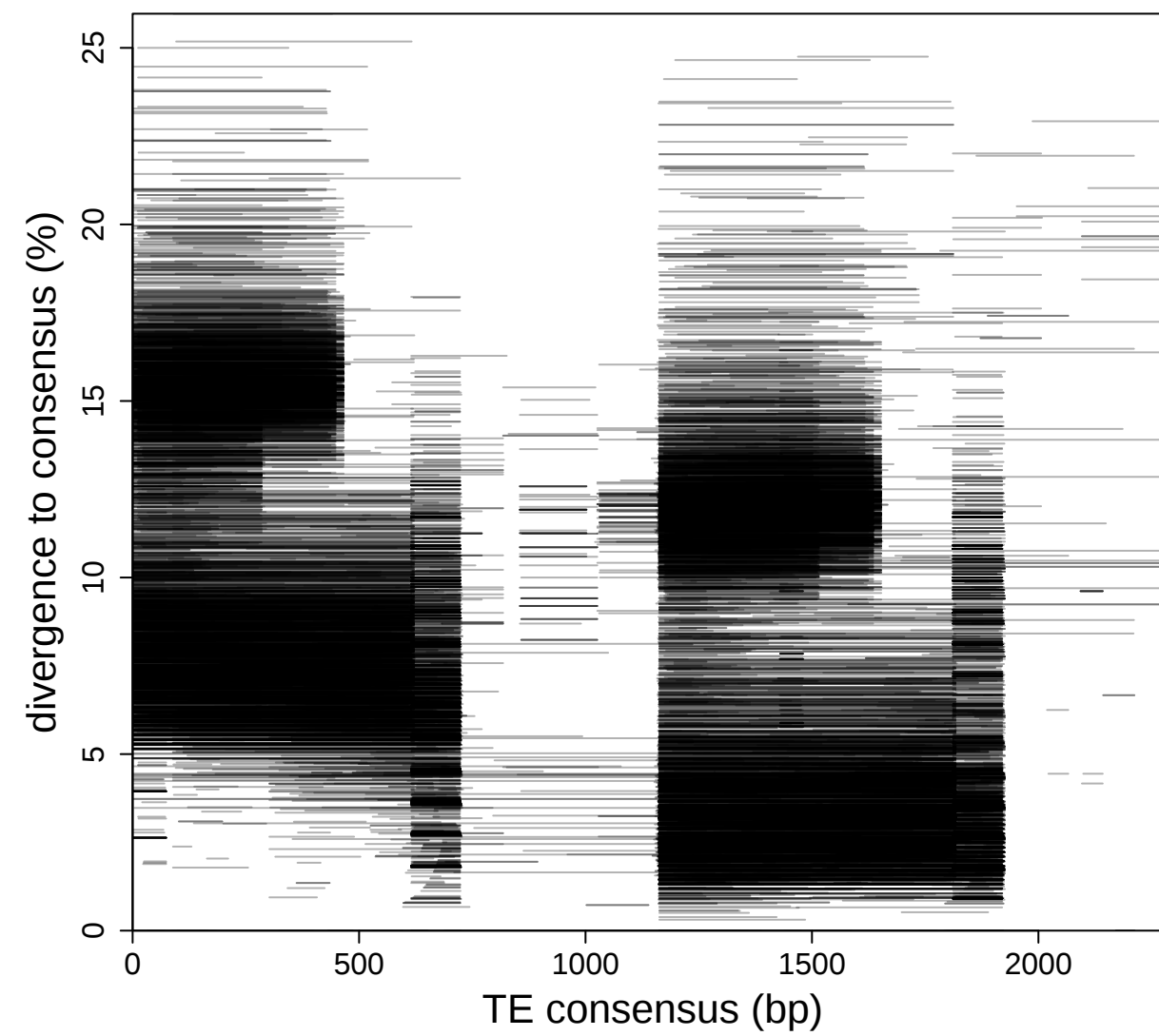
Start crop Point

End crop Point

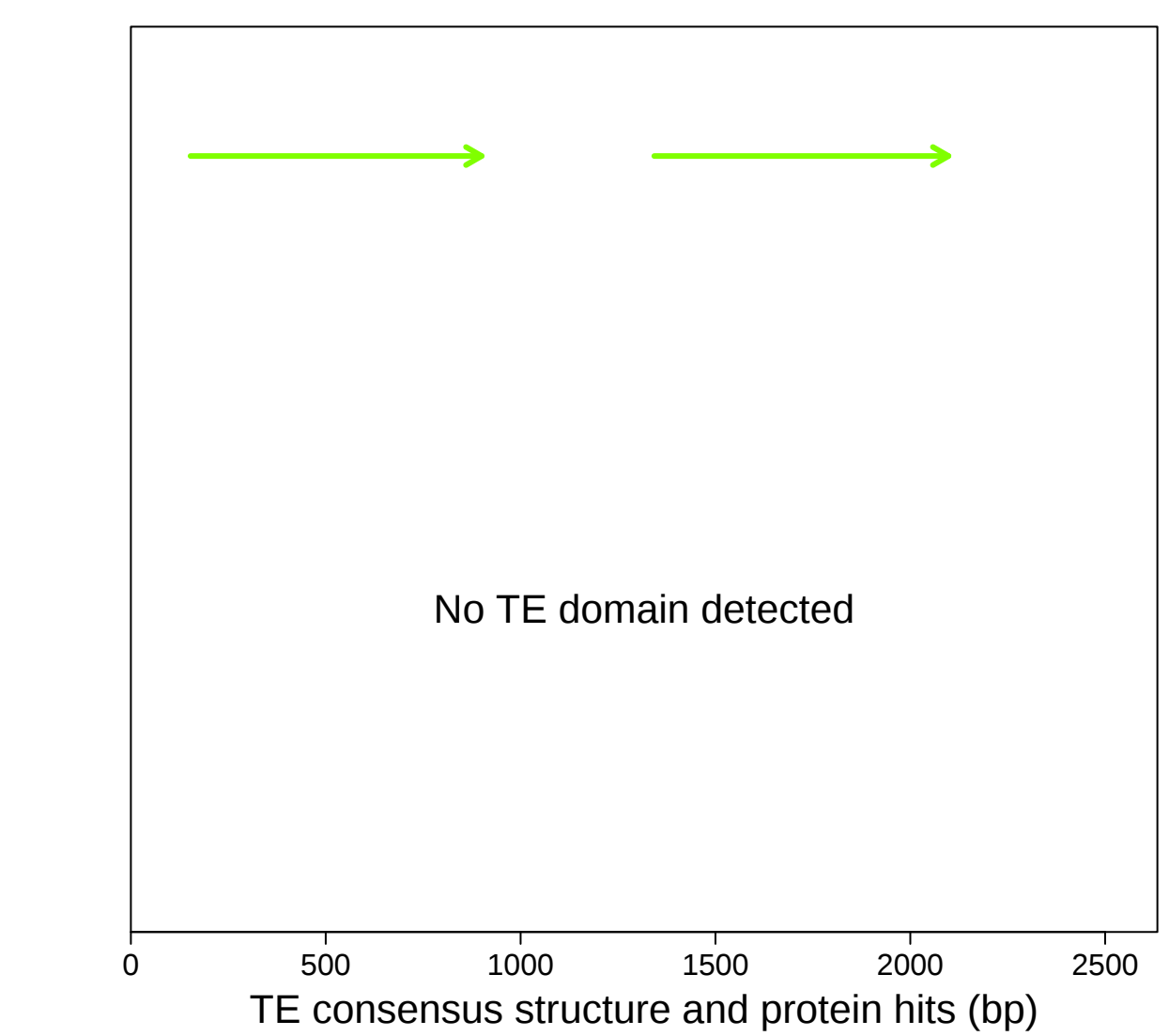
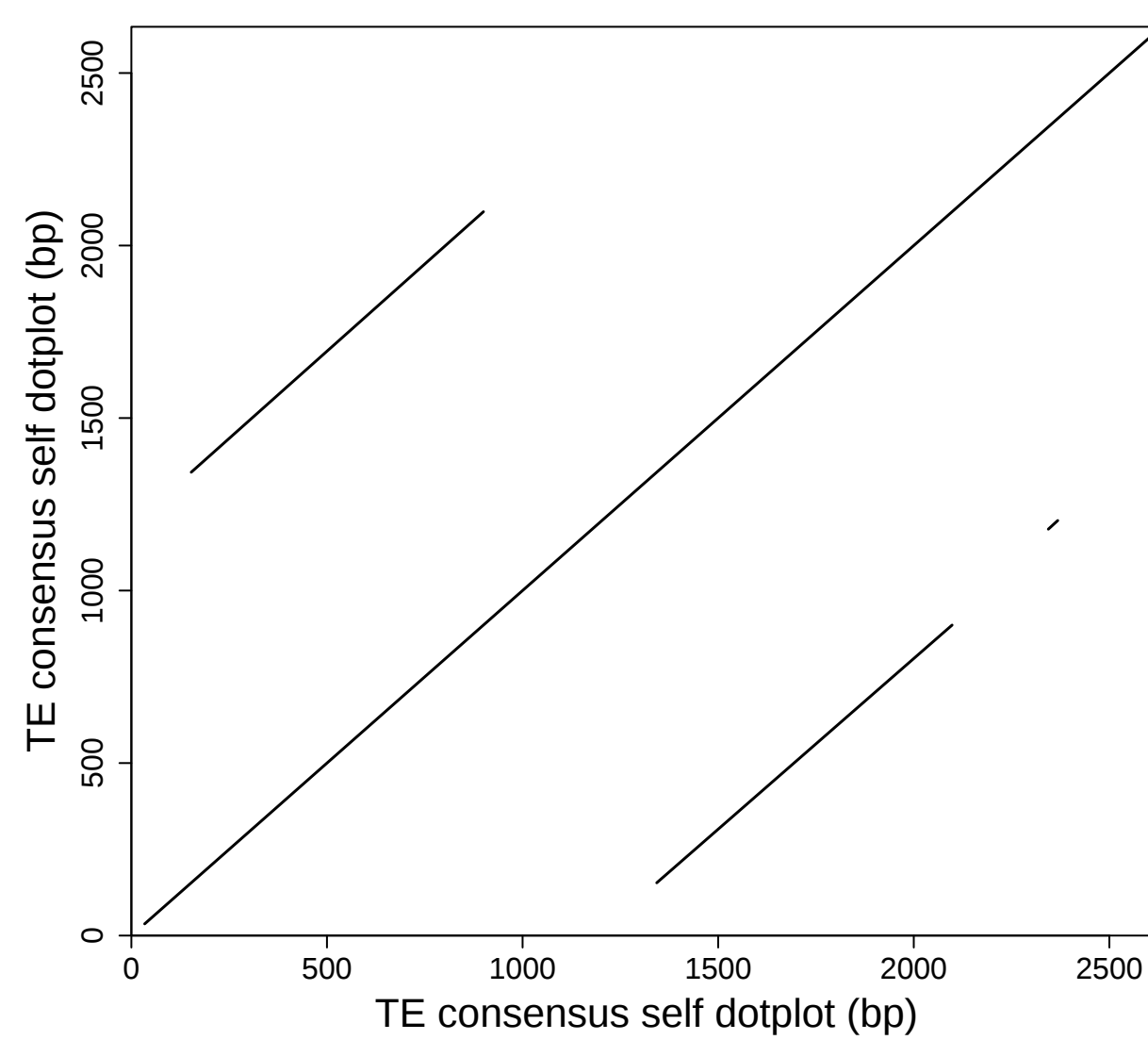
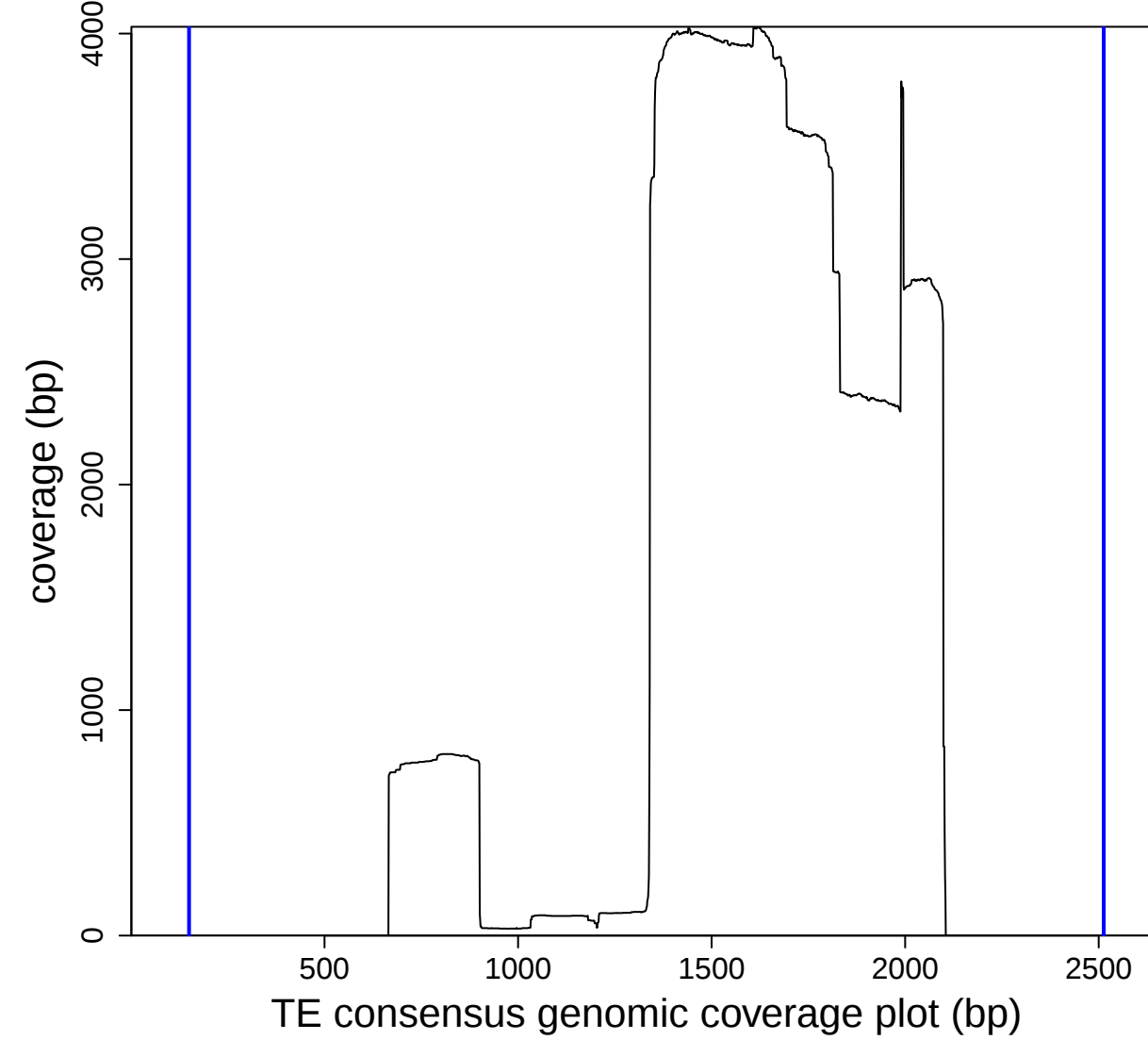
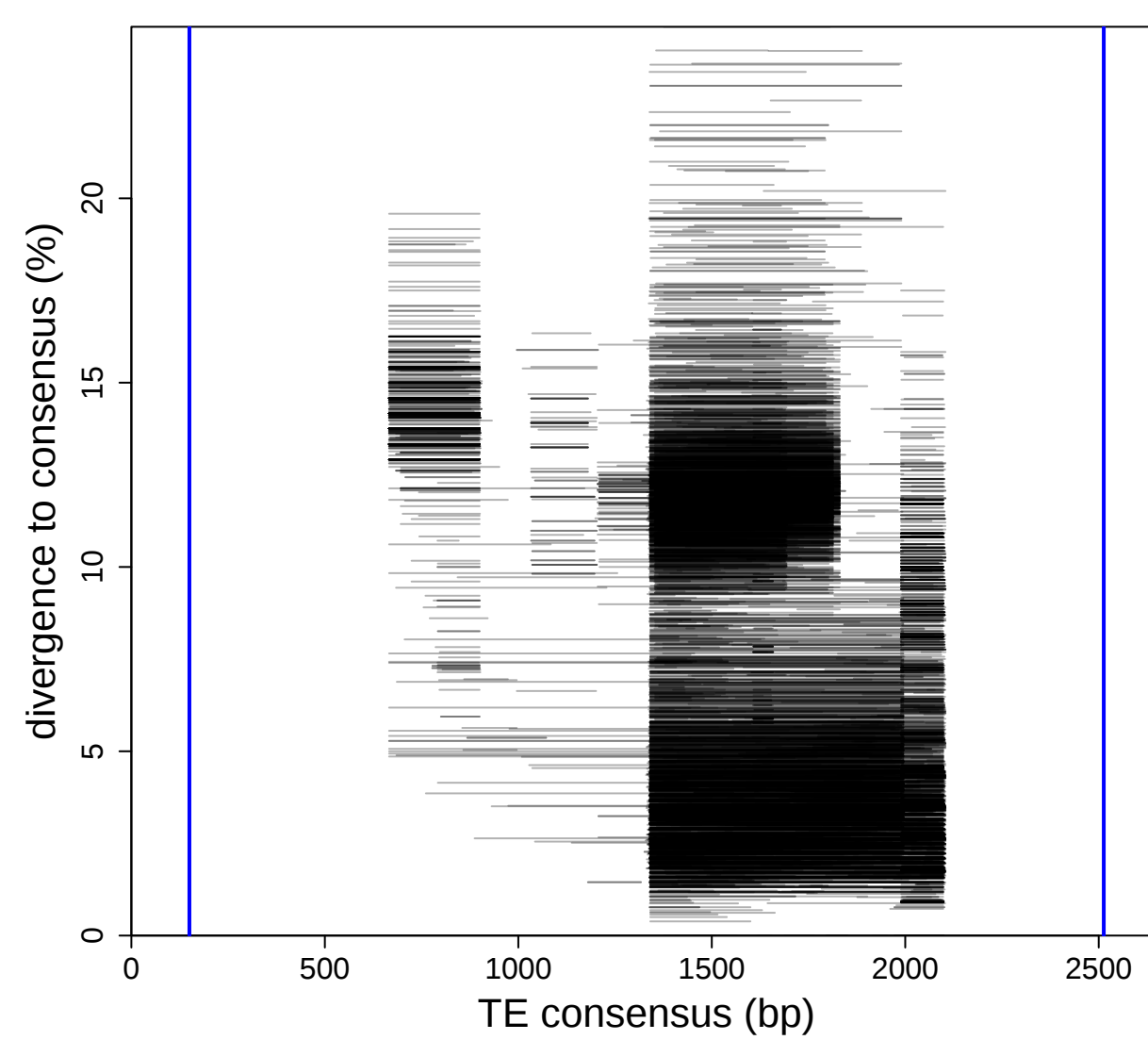


bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.fa
size: 2297bp; fragments: 12831; full length: 0 (>=2067.3bp)

After TEtrimmer 2297 bp



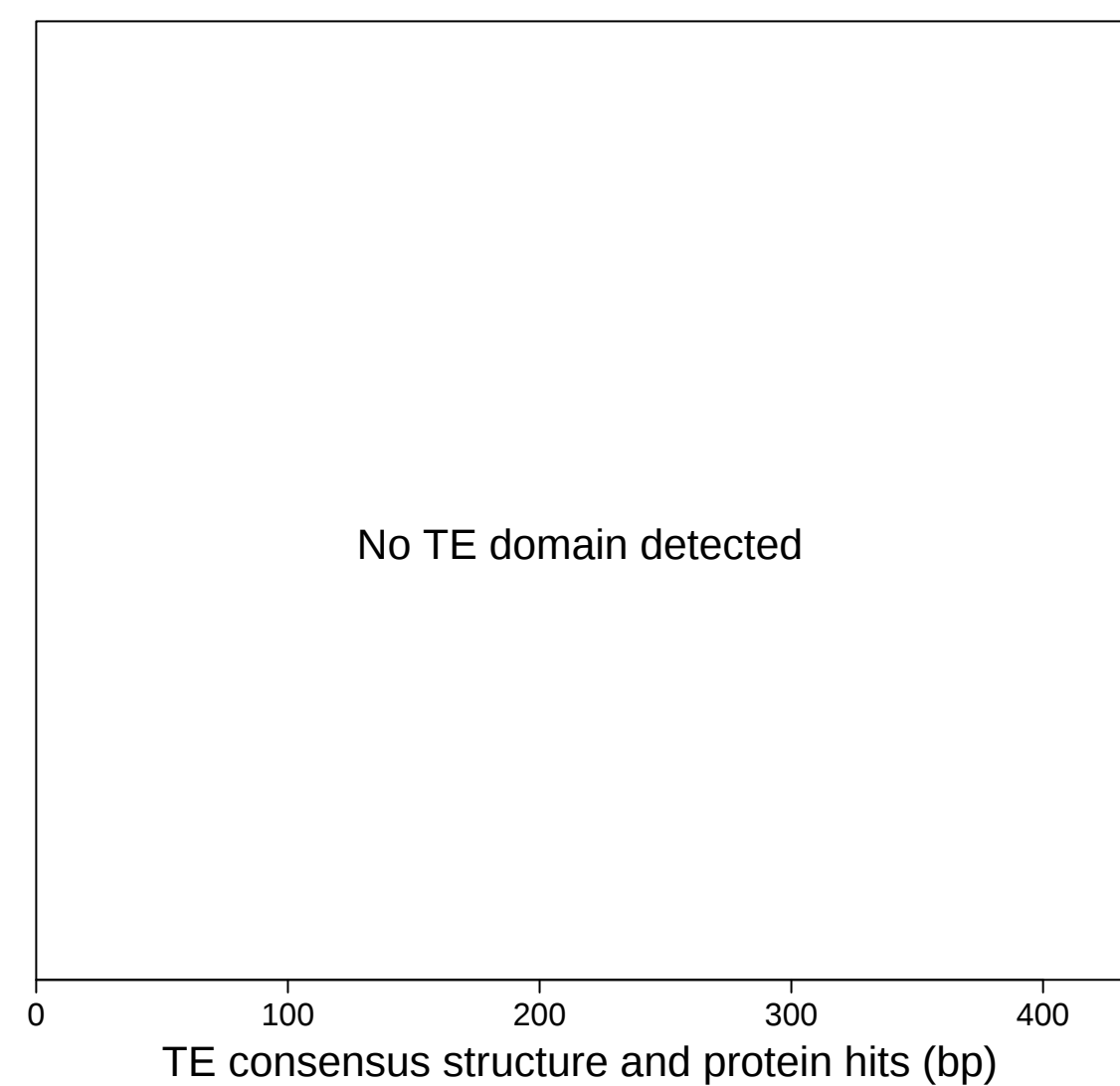
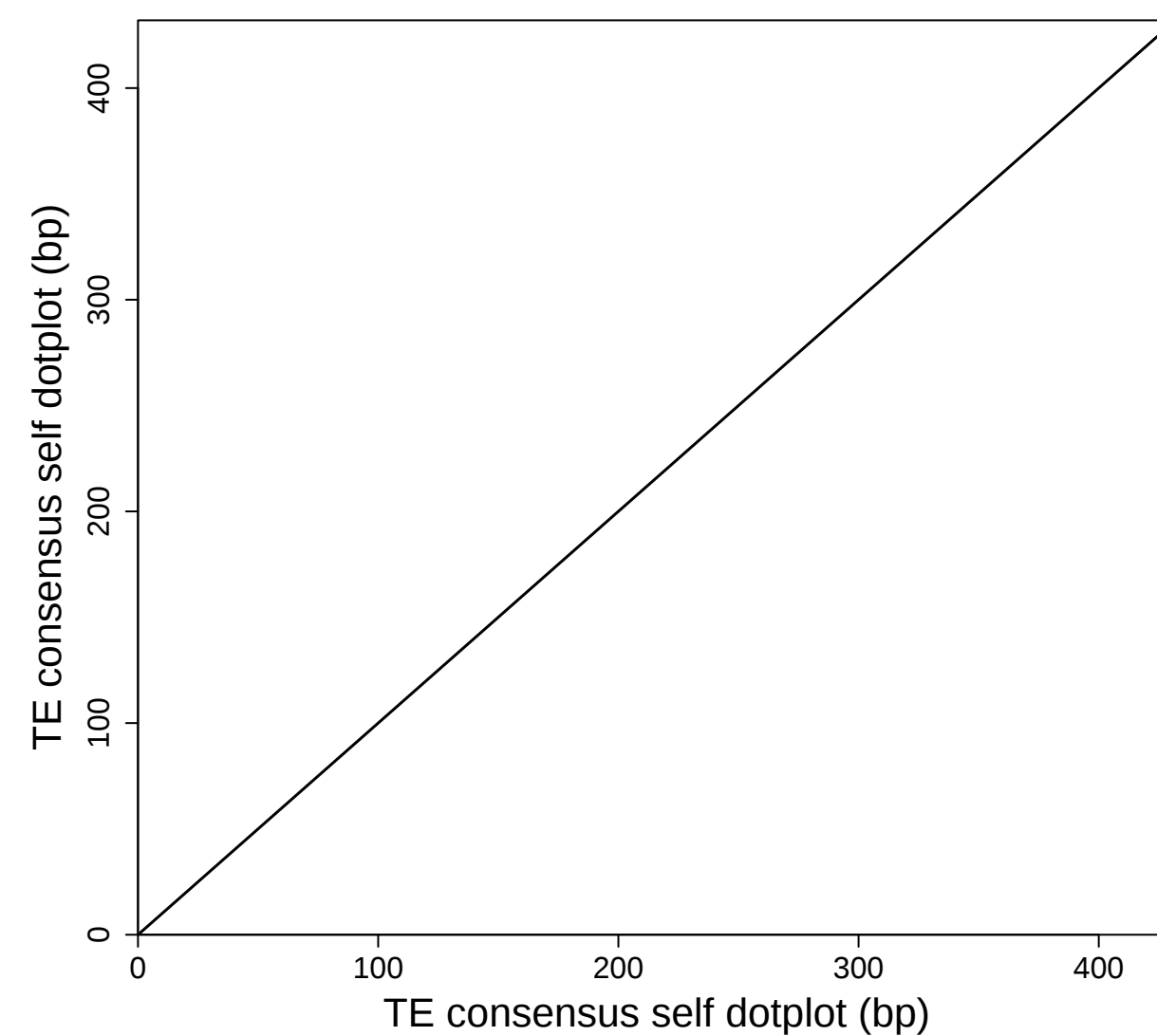
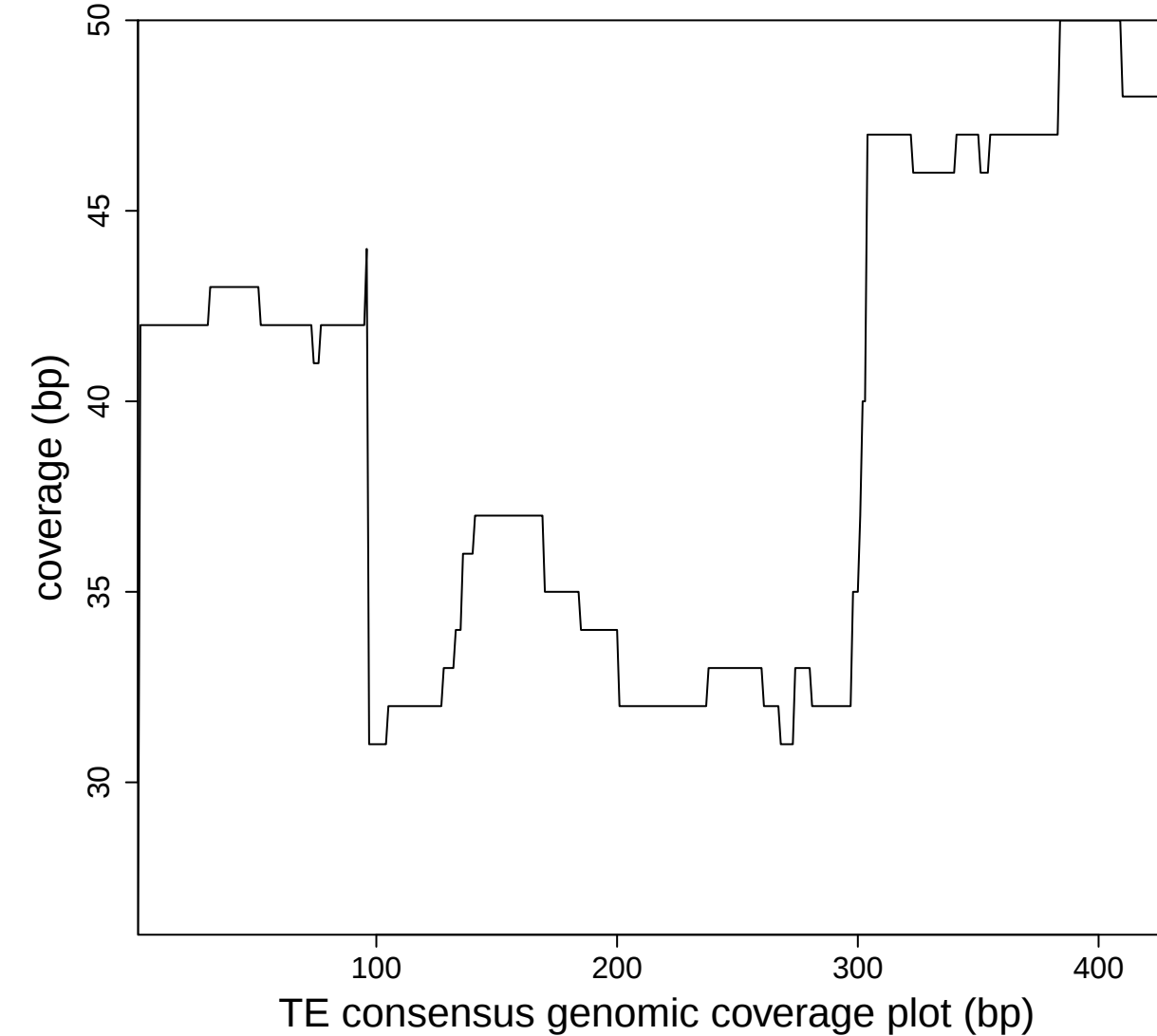
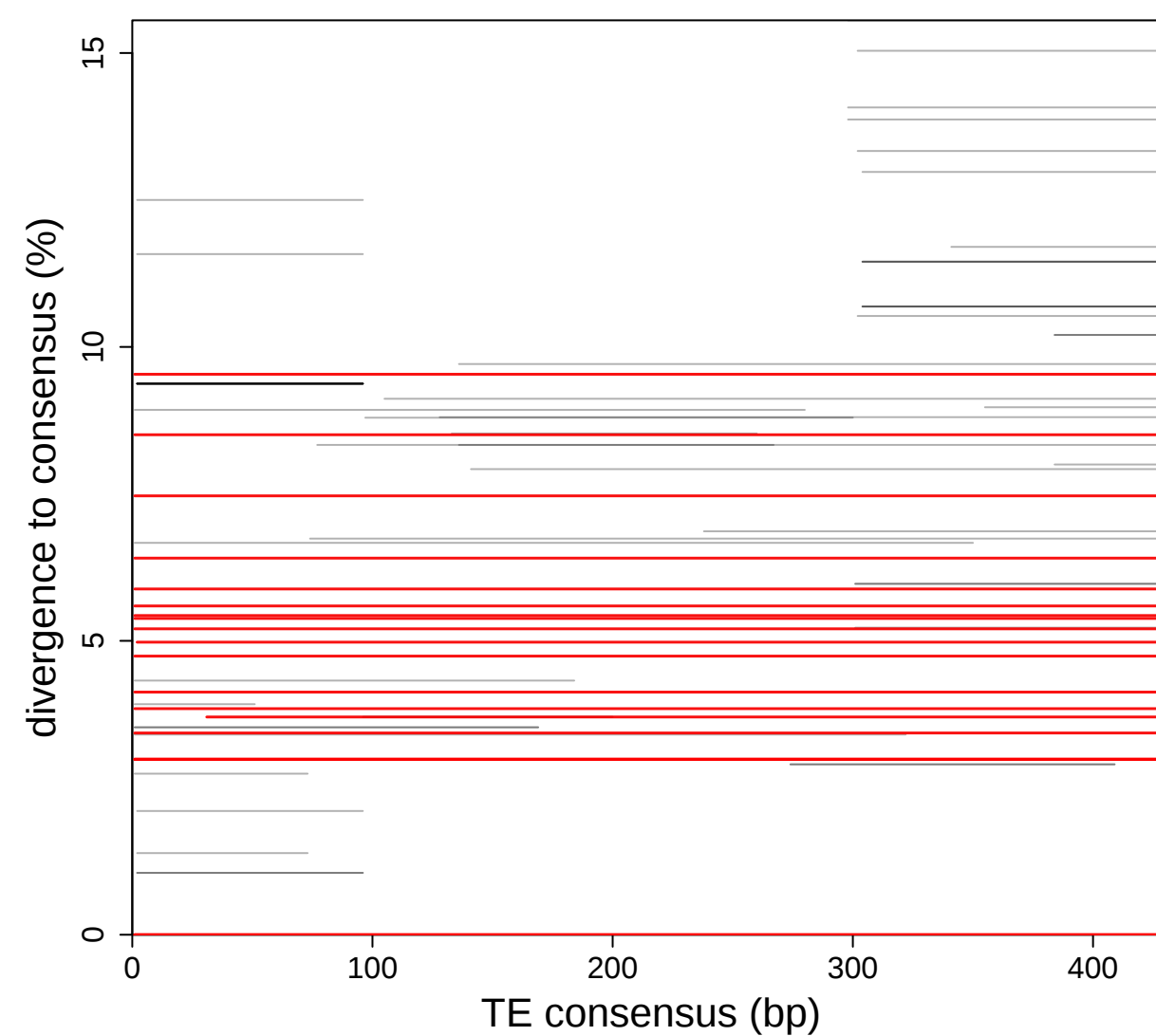
ed_uf.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_
size: 2663bp; fragments: 7375; full length: 0 (>=2396.7bp)



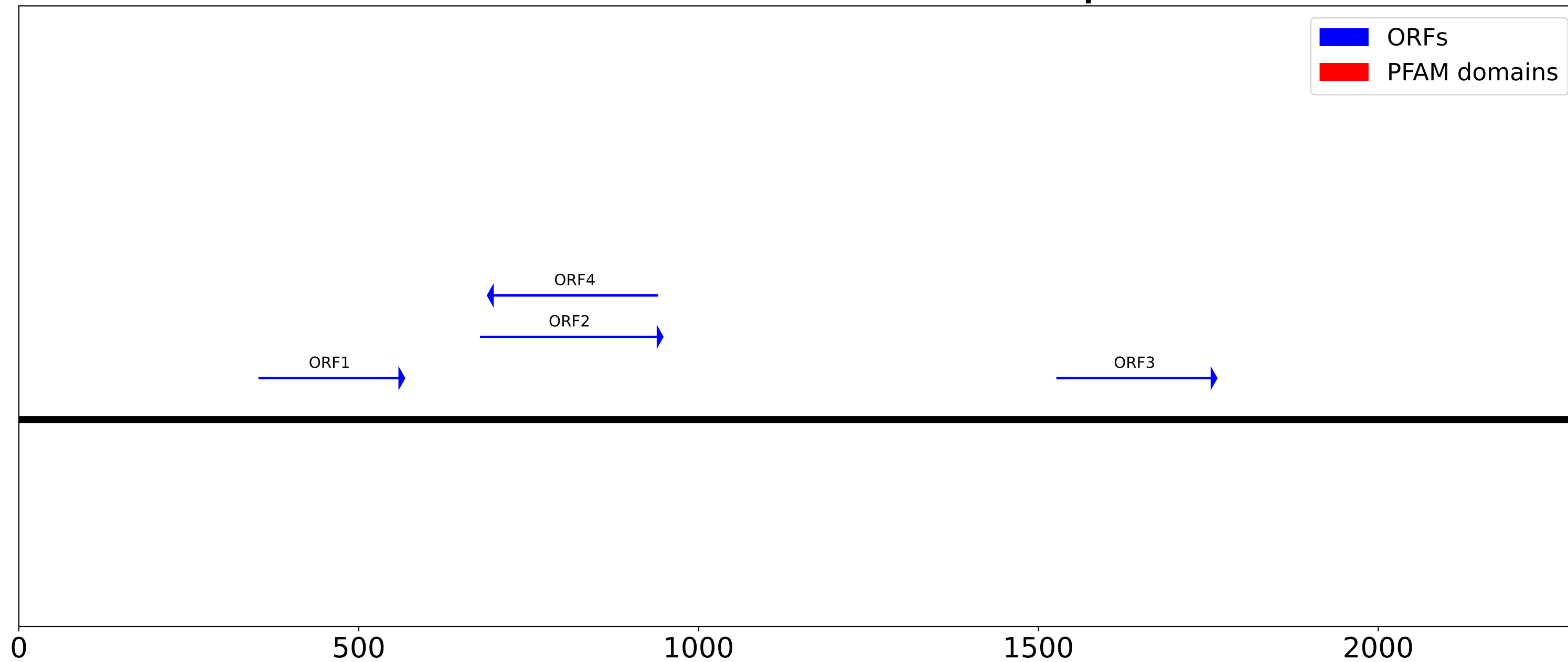
Before TEtrimmer 432 bp

TE: ltr_1_family_758

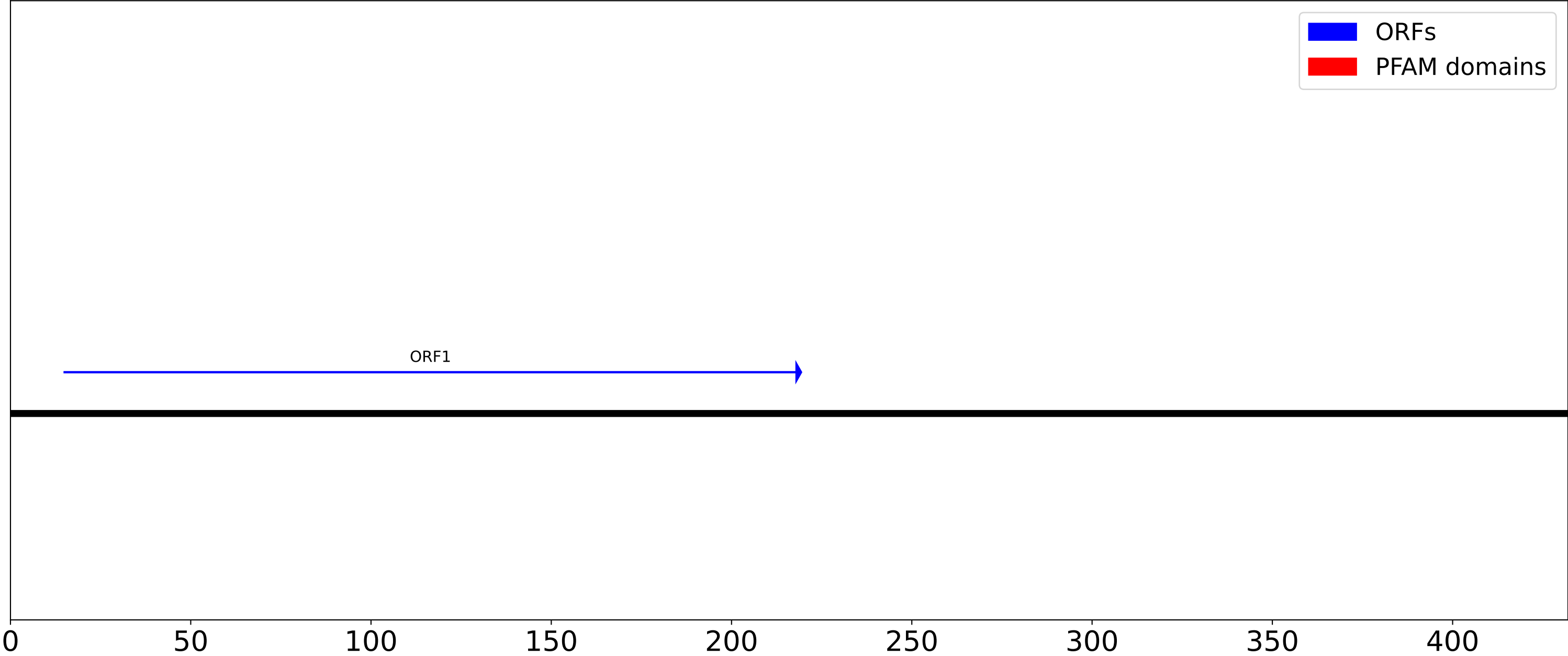
size: 432bp; fragments: 78; full length: 20 (≥ 388.8 bp)



After TEtrimmer ORF and PFAM domain plot



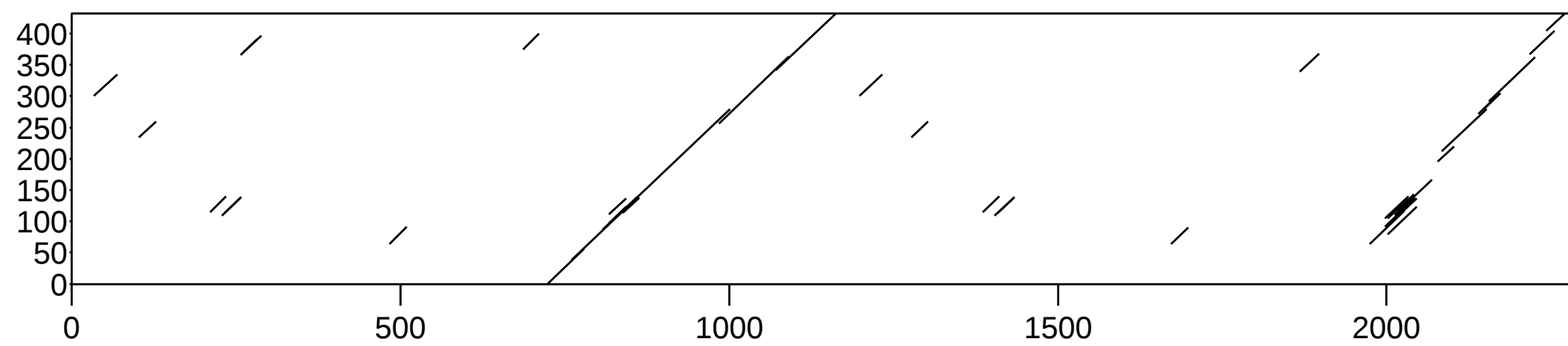
Before TEtrimmer ORF and PFAM domain plot



Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)